4. (Once amended) A transformed plant having a nucleic acid molecule which

comprises:

(a)

(a) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule;

- (b) a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 and complement thereof;
- (c) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.
- 5. (Once amended) The transformed plant according to claim 4, wherein said nucleic acid molecule comprises a complement of a nucleic acid sequence SEQ ID NO: 1.

8. (Once amended) A method for determining a level or pattern in a plant cell or plant tissue of a protein in a plant comprising:

incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 or complement thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid

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Sub contid (c)

molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said protein; permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said protein.